



Figure S1. Crystal structure of HIV-1 Tat complexed with ATP-bound human P-TEFb. Residues 1 to 48 of Tat protein are resolved in the structure with visible α -helices for the C-terminus region (residues 28-43). Despite unstructured, the N-terminus of Tat protein is stabilized by the P-TEFb protein in the complex. Cyclin T1 is colored in dark green, CDK9 in grey and Tat in purple. This figure was prepared with data from PDB entry 3MIA.